

Hale, Mary

160141

From: Angell, Jon E  
Sent: Wednesday, July 27, 2005 11:31 AM  
To: Hale, Mary  
Subject: RE: 09/899,276

Hello Mary,

I just finished talking to Jeff Fredman and he recommended requesting the following search:

A standard search performed using the following SEQ. ID NOs. from application : 09/899,276

SEQ ID NO: 1 (nucleic acid 600 nucleotides long)  
SEQ ID NO: 2 (nucleic acid 200 nucleotides long)  
SEQ ID NO: 3 (nucleic acid 150 nucleotides long)  
SEQ ID NO: 4 (nucleic acid 250 nucleotides long)  
SEQ ID NO: 5 (nucleic acid 300 nucleotides long)  
SEQ ID NO: 6 (nucleic acid 300 nucleotides long)  
SEQ ID NO: 8 (nucleic acid 20 nucleotides long)

AND, if possible, nucleotides 8968-9073 of SEQ ID NO: 13 followed by approx. 50 Ns (N=any nucleotide) followed by the sequence TGATCA.

Please note that the previous search request indicated that standard and oligo searches were required, that is no longer the case. Only a standard search is required.

(note: previous search request is pasted below, for your convenience)

Thanks,

Eric

-----Previous search request below -----

Examiner#: 78697  
Art Unit : 1635  
Phone Number: 571-272-0756  
Date: 7/21/2005  
Serial Number: 09/899,276  
Mailbox & Bldg/Room Location: REMSEN 2C18  
Results Format Preferred (circle): Paper

VOID See  
above modification/  
revision.  
W. Hale  
7/26/05

I would like to have a standard search performed using the following SEQ. ID NOs. from application : 09/899,276

SEQ ID NO: 1 (nucleic acid 600 nucleotides long)  
SEQ ID NO: 2 (nucleic acid 200 nucleotides long)  
SEQ ID NO: 3 (nucleic acid 150 nucleotides long)  
SEQ ID NO: 4 (nucleic acid 250 nucleotides long)  
SEQ ID NO: 5 (nucleic acid 300 nucleotides long)  
SEQ ID NO: 6 (nucleic acid 300 nucleotides long)  
SEQ ID NO: 8 (nucleic acid 20 nucleotides long)

SEQ ID NO: 13 (nucleic acid 11793 nucleotides long)

Please note that for SEQ ID NOS: 1, 2, 3, 4, 5, 6 and 8 **standard and oligo** searches of the nucleic acid databases are required.

For SEQ ID NO: 13, a **standard** search of the nucleic acid database and a **cross-search** in the protein databases (to

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identify protein encoded by SEQ ID NO 13) are required.

-----Original Message-----

**From:** Hale, Mary  
**Sent:** Wednesday, July 27, 2005 9:35 AM  
**To:** Angell, Jon E  
**Subject:** 09/899,276

Hi Jon,

The sequence search submitted on 7/21/05, sequence id #13 has 11,793 residues which is over the 10,000 residue limit. Sequence id #13 has been canceled but all the other sequences will be searched. Please meet with one of the examiner screen team members (Martinell, Pak, Priebe, Fredman). They will review your request and offer suggestions on optimizing the processing time.

I will hold on to your request until noon today and if you are able to meet and modify the sequence, we will process sequence id 13 with the other sequences. Please e-mail me directly with your changes. If I do not hear from you by noon today, seq id 13 will not be included in the search.

If you have any questions, please contact me.

Mary Hale  
2-2507

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